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Command line parameters:

-MODEL-frame+_p2n.model -DEV=xlh
-Q-/cgn2_1/USPTO_spool/US09807459/runat_18102002_141112_28595/app_query.fasta_1.647
-Q-/cgn2_1/USPTO_spool/US09807459/runat_18102002_141112_28595/app_query.fasta_1.647
-Q-/cgn2_1/USPTO_spool/US09807459/runat_18102002_141112_28595/app_query.fasta_1.647
-DB-EST -OFMT=fastap -SUFFIX=rst -MINMATCH=0.1 -LOOPCL=0 -LOOPEXT=0
-UNITS=bits -START=1 -END=-1 -MATRIX=Dlosum62 -TRANS=Duman40.cd1 -LIST=45
-DOCALIGN=200 -THR_SCORE-pct -THR_MAX=100 -THR_MIN=0 -ALIGN=40 -MODE=LOCAL
-OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=200000000
-USER-US09807459_eCGN_1_1763_erunat_18102002_141112_28595 -NCPU=6 -ICPU=3
-NO_XLPXY -NO_MMAP -LARGEQUERY -NEG_SCORES=0 -WAIT -LONGLOG -DEV_TIMEOUT=120
-WARN_TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPOEXT=0.5 -FGAPOP=6 -FGAPEXT=7
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CH230-48P RPCI-24-3	1M0199K2	CH230	1M0075E05	CH230-101	1M0465N20		RPCI-24-2	ENTKT09T	RPCI-24-2	ENTFZ06TR	RPCI-23-2	1M0078M05	2M0208J13	RPCI-24-2	31-25 Ste	RPCI-24-2	EST570974	2M0205C06	PfESToaa6	1M0271A02	RPCI-24-1	1M0216006	T3 end of	T3 end of	2M0082P07	Mus musc	BB230024	2M0271021	SP_0034_B	1M0564C11	2 RPCI-24-2		Tetrac	1M0167E11	SP_103	SP_0001_	ש	PfESToaa	OP21695	SP_1039	SP_0106_	

ALIGNMENTS

Cameron,R.A., Mahairas,G., Rast,J.P., Martinez,P., Biondi,T.R., Swartzell,S., Wallace,J.C., Poustka,A.J., Livingston,B.T., Wray,G.A., Ettensohn,C.A., Lehrach,H., Britten,R.J, Davidson,E.H. ar AZ167872 529 bp DNA linear SP_0103_AZ=E10_T7A Strongylocentrotus purpuratus, sperm genomic BAC library Strongylocentrotus pur clone Plate=103 Col=20 Row=I, DNA sequence. Contact: Cameron, RA, Davidson, EH, Division of Biology 156-29 A sea urchin genome project: Sequence scan, virtual map, and additional resources Strongylocentrotus purpuratus Eukaryota; Metazoa; Echinodermata; Eleutherozoa; Echinozoa; Echinoidea; Euechinoidea; Echinacea; Echinoida; GSS AZ167872 AZ167872.1 GI:8338240 California Institute of Technology 20402566 Hood, L. Strongylocentrotus purpuratus. Strongylocentrotidae; Strongylocentrotus (bases 1 to 529) Natl. Acad. Sci. U. S. A. 97 (17), Hood, L 9514-9518 (2000) purpuratus genomic GSS 29-AUG-2000 purple sea urchin

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RESULT 2
BM275292
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Tel: (626) 395-8421
Fax: (626) 793-3047
Email: acameron@caltech.edu
Plate: 103 row: I column: 2
Seq primer: T7
Class: BAC ends
                                     BM275292 564 bp mRNA linear EST 20-DE PfESToaa66f11.yl Plasmodium falciparum 3D7 gametocyte cDNA 1i Plasmodium falciparum 3D7 cDNA 5' similar to TR:097255 097255
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                      PROTEIN.
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DH10B"
1 155 c 158 g 94 t
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/clone_lib="Strongylocentrotus purpuratus,
urchin, sperm genomic BAC library"
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/db_xref="taxon:7668"
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US-09-807-459-2 (1-458) x BM275292 (1-564)
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{\tt AspAlaLysAspLeuPheGluAsnLysIleGlyGlnGlyThrValAspPheIleAsnAsn}
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Fax: 314 286 1810
Email: estewatson.wustl.edu
Library was constructed by R. Haywood. DNA sequencing
Washington University Genome Sequencing Center For inf
obtaining a clone please contact: L. David Sibley
(sibley@borcim.wustl.edu), Washington University
Seq primer: -40Up from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WashU Plasmodium EST Project
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108, USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Tang,K., Cole,R., Chakrabarti,D., Haywood,R., Clifton,S., Pape,D., Marra,M., Hillier,L., Martin,J., Wylie,T., Dante,M., Theising,B., Bowers,Y., Gibbons,M., Ritter,E., Bennett,J., Jentes,E., Ronko,I., Tsagareishvili,R., Belaygorod,L., Franklin,C., Carr,L., Grow,A., Maguire,L., Richey,J., Wadkins,J., Kennedy,S., Levinso,D., Waterston,R., Wilson,R. and Sibley,D.
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1 (bases 1 to 564)
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Plasmodium falciparum 3D7
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Location/Qualifiers
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/lab host="ph10B (GeneHog, Invitrogen, Inc.)"
/lab host="ph10B (GeneHog, Invitrogen, Inc.)"
/note="Wector: pBluescript SK plus; Site_1: EcoRI; Site_2: XhoI; The library was constructed by R Haywood. cDNAs were synthesized from ganetocyte poly(A)+ RNA by oligo d(T) priming, size-selected and directionally cloned into the EcoRI (5' end) to XhoI (3' end) sites of the Uni-ZAP XR lambda vector (Stratagene). The primary library was mass excised as phagemid using the ExAssist helper phage (Stratagene) clones were mass excised using the ExAssist helper phage (Stratagene), the phagemids were precitptated with PEG 8000 and extracted with phenolychloroform. Phagemid DNA was electroporated into DH10B cells. Clone Availability: David Sibley, Washington University."
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952 bp DNA linear GSP_0106_A2_A12_SP6E Strongylocentrotus purpuratus, pu urchin, sperm genomic BAC library Strongylocentrotus genomic clone Plate=106 Col=24 Row=A, DNA sequence.
AZ209422
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California Institute of Technology
Pasadena California 91125, USA
Tel: (626) 395-8421
Fax: (626) 793-3047
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Eukaryota; Metazoa; Echinodermata; Eleutherozoa; Echinozoa;
Echinoidea; Euechinoidea; Echinacea; Echinoida;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: acameron@caltech.edu
plate: 106 row: A column:
Seq primer: SP6
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/clone_lib="Strongylocentrotus purpuratus, purple sea
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Eukaryota; Metazoa; Echinodermata; Eleutherozoa;
Echinoidea; Euchinoidea; Echinacea; Echinoida;
Strongylocentrotidae; Strongylocentrotus.

1 (bases 1 to 856)
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SP_1039_B2_E12_T7A Strongylocentrotus purpuratus, purple sea urchin, sperm genomic BAC library Strongylocentrotus purpuratus genomic clone Plate=1039 Col=24 Row=J, DNA sequence.
                                                                                                                                                                                                                                                                                                                                 Division of Biology 156-29
California Institute of Technology
Pasadena California 91125, USA
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                                                                                                                                                             Class: BAC ends
                                                                                                                                                                                                                         Email: acameron@caltech.edu
Plate: 1039 row: J column:
                                                                                                                                                                                                                                                                              Tel: (626) 395-8421
Fax: (626) 793-3047
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Az199582.1 GI:8394390
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Location/Qualifiers
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                    /organism="Strongylocentrotus purpuratus"
/db_xref="taxon:7668"
/clone="Plate=1039
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BM416596
DP216596
DP216595
Mixed Stage EST's from Globodera pallida, the potato cyst nematode Globodera pallida cDNA similar to Homology: (blast score <-3D 1e-5) Drosophila melanogaster genomic scaffold 142000013386053 section 27 of 30, mRNA sequence.

BM416596.1 GI:18383396
                                                                                                                                                                     Globodera pallida.
Globodera pallida
Eukaryota, Metazoa, Nematoda; Chromadorea; Tylenchida;
Tylenchoidea; Heteroderidae; Heteroderinae; Globodera.
Contact: Opperman, C
Center for the Biology of Nematode
Center for the Biology of Nematode
NC State University; IACR-Rothamste
Campus Box 7616; Raleigh, NC 27695,
Tel: 919.515.6699
Fax: 919.515.9500
                                                                                                      Heer, J., Sosinski, B., Pokrzywa, R.M., Warry, A. Mixed Stage EST's from Globodera pallida, the Unpublished (2001)
                                                                                                                                                           (bases 1 to 1035)
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PCNGT11-3_R_F04_PCN_R_031.ab1
Location/Qualifiers
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/db_xref="taxon:36090"
/clone_lib="Mixed Stage EST's from Globodera
potato cyst nematode"
/note=""""
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note-"Vector: lambda GT11; This is a collaborative efformetween IACR-Rothamsted and North Carolina State University. The library was constructed from mixed stage G. pallida in lambda GT11 by Paul Burroughs, IACR-Rothamsted."
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PhePheAsnLysGluIleArgAspPro---SerLysAlaLeuLysGluLysValSerAsn 345
                                                   CCAATAAATGACCCAATAAATGACCCAATAAATGACCCAATAAATGACCCAATAAATGAC
                                                                                                    LysValSerThrAspThrLysAspLeuPheGluAsnLysIleGlyGlnGlyThrValAsp 326
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Email: est@watson.wustl.edu
Library was constructed by R. Haywood. DNA sequencing
Washington University Genome Sequencing Center For inf
obtaining a clone please contact: L. David Sibley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WashU Plasmodium EST Project washington University School of Medicine 4444 Forest Park Parkway, Box 8501, St. I Tel: 314 286 1800
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Location/Qualifiers
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//lab_host="DHIOB (GeneHog, Invitrogen, Inc.)"
//note="Vector: pBluescript SK plus; Site_1: EcoRI; Site_2: XhoI: The library was constructed by R Haywood. cDNAs were synthesized from ganetocyte poly(A)+ RNA by oligo d(T) priming, Size-selected and directionally cloned into the EcoRi (5' end) to XhoI (3' end) sites of the Uni-ZAP XR lambda vector (Stratagene). The primary library was mass excised as phagemid using the Exassist helper phage (Stratagene), the phagemids were precitytated with PEG 8000 and extracted with phenol/chloroform. Phagemid DNA was electroporated into DHIOB cells. Clone Availability: David Sibley, Washington University."

53 a 130 c 42 g 78 t 1 others
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WashU Plasmodium EST Project
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BM274676 537 bp mRNA linear EST 20-DEC-200 PfESToaa73c03.yl Plasmodium falciparum 3D7 gametocyte cDNA library Plasmodium falciparum 3D7 cDNA 5' similar to TR:097255 097255
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Fax: 314 286 1810
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Plasmodium falciparum 3D7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: est@watson.wustl.edu
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XhoI; The library was constructed by R Haywood. cDNAs were synthesized from ganetocyte poly(A) + RNA by oligo d(T) priming, size-selected and directionally cloned into the EcoRi (5' end) to XhoI (3' end) sites of the Uni-ZAP XR lambda vector (Stratagene). The primary library was mass excised as phagenid using the Exassist helper phage (Stratagene). Clones were mass excised using the Exassist
                                                                                                                                                                                                                                                                       /organism="Plasmodium falciparum 3D7"
/db_xref="taxon:36329"
/clone_lib="Plasmodium falciparum 3D7
                                                                                                                                                                      /dev_stage="gametocyte (stage III-V)"
/lab_host="DH10B (GeneHog, Invitrogen, Inc.)"
/note="Vector: pBluescript SK plus; Site_1: EcoRI; Site_2://opte="Vector"
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                                                                                                                                               Strongylocentrotus purpuratus
Eukaryota; Metazoa; Echinodermata; Eleutherozoa;
Echinoidea; Euechinoidea; Echinacea; Echinoida;
Strongylocentrotidae; Strongylocentrotus.
                                             A sea urchin genome project: Sequence scan, virtual map, additional resources
                                                                                                     Cameron,R.A., Mahairas,G., Rast,J.P., Martinez,P., Biondi,T.R., Swartzell,S., Wallace,J.C., Pouska,A.J., Livingston,B.T., Wray
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                                                                                         AZ199581
SP 1039 B2_E12_SP6E Strongylocentrotus purpuratus, purple sea urchin, sperm genomic BAC library Strongylocentrotus purpuratus genomic clone Plate=1039 Col-24 Row-J, DNA sequence.
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California Institute of Technology
Pasadena California 91125, USA
Strongylocentrotus purpuratus
Eukaryota; Metazoa; Echinodermata; Eleutherozoa;
Echinoidea; Euechinoidea; Echinacea; Echinoida;
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Class: BAC ends
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Fax: (626) 793-3047
                                             GSS.
Strongylocentrotus purpuratus.
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/clone="Plate=1 Col=2 Row=N"
/clone_lib="Strongylocentrotus purpuratus,
urchin, sperm genomic BAC library"
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AZ400717
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California Institute of Technology
Pasadena California 91125, USA
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Class: BAC ends
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Plate: 1039 row: J column: 24
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Fax: (626) 793-3047
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urchin, sperm genomic BAC library"
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Insert Length: 10000 Std Erro
Plate: 0167 row: E column: 1
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University of Utah
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                                                                                                                                                                                                                                                                                                                                                                               was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWDA2 (gi14732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the coort adaptors and
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone="UUGC1M0167E11"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
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/strain="C57BL/6J"
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GSS; genome survey sequence.
GSS: genome nigroviridis.
Tetraodon nigroviridis
Tetraodon nigroviridis
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontidae; Tetraodon.
Bernot, A., Fiz
Saurin, W. and
           Roest-Crollius, H., Jaillon, O., Dasilva, C., Bouneau, L., Fisher, C
Bernot, A., Fizames, C., Wincker, P., Brottier, P., Quetler, F.,
                                       Unpublished
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                                                                 Charaterization and repeat analysis of the compact genome freshwater pufferfish Tetraodon nigroviridis
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Roest-Crollius, H., Jaillon, O., Dasilva, C., Fizames, C., Fizames, L., Billault, A., Quetier, F., Saurin, W., Bernot, A.
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                                                                                                                                                                                                                                                                            Contact: Marra M/Mouse EST Project Washington University School of MedicineP 4444 Forest Parkway, Box 8501, St. Ld Tel: 314 286 1800

Fax: 314 286 1810
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                                                                                                                                                                                                                                              Email: mouseest@watson.wustl.edu
This clone is available royalty-free through LLNL ;
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                                                                                                                                                                                                                      IMAGE Consortium (info@image.llnl.gov)
MGI:561948
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ·CTTTCAGAATCAGAGAWCATTAAGGGGAATGTYGATTTTCTCATCAGAGAGAAAAG
                                                                                                                                                                                        quality sequence stop: 469 Location/Qualifiers
 //note="Organ: embryo; Vector: pSPORT; Site_1: NotI;
Site_2: SalI; Cloned unidirectionally from mRNA prepared
from 800 blastocysts. primer: SalI(dT):
5'.CGGTCGACCGTCGACCGTTTTTTTTTTTTTTT-3'. cDNAs were
                                                        /clone_lib-"Knowles Solter mouse blas
/tissue_type-"blastccyst"
/dev_stage-"embryo (pre-implantation)
/lab_host-"DH10B"
                                                                                                                 /organism="Mus musculus"
/strain="B6D2 F1/J"
/db_xref="taxon:10090"
/clone="IMAGE:989668"
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Contact: Shaying Zhao
Department of Eukaryotic Genomics
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US-09-807-459-2 (1-458) x AA571652
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Unpublished (1999)
Other_GSSs: RPCI-24-287H12.TJ
                                 Zhao, S., Nierman, W., Malek, J., Shatsman, S., A
Tsegaye, G., Geer, K., Krol, M., Shvartsbeyn, A.,
Russell, D., de Jong, P. and Fraser, C.M.
Mouse BAC End Sequences from Library RPCI-24
                                                                                                           Mus musculus
Eukaryota; Metazoa; Chordata;
Eukharia; Rodentia;
1 (bases 1 to 794)
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----GlyThrValAsp---PheIleAsnAsnGluIleArgAspProSerLysAlaLeuIl 375
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2 Medical Center Dr., Rockville, MD 20850, USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /db_xref="taxon:10090"
/clone="RPCI-24-287H12"
/clone_lib="RPCI-24"
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/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0564 row: C column: 11
Seg primer: CACACAGGAAACAGCTATGACC
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AZ766807

IM0564C11R Mouse 10kb plasmid UUGC1M library Mus musculus genomic Clone UUGC1M0564C11 R, DNA sequence.
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  adaptored DNA was purified and size-selected for a 9.5 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derival
                                                        was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                              (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA
                                                                                                                                                                                            /lab_host="E. Coli strain XL10-Gold, Tl-resistant, F
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jack
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electrophoresis.
                                                                                                                                                                            Laboratory Mouse DNA Resource
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/clone="UUGCIM0564C11"
/clone_11b="Mouse 10kb plasmid UUGCIM library"
                                                                                                                                                                                                                                                                                                                                   /organism="Mus musculus"
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Strongylocentrotus purpuratus.
Strongylocentrotus purpuratus.
Strongylocentrotus purpuratus
Eukaryota; Metazoa; Echinodermata; Eleutherozoa;
Eukaryota; Metazoa; Echinoida; Echinoida;
                                                                                                                         AZ152840 324 bp DNA linear GSP_0034_B2_G06_SP6E Strongylocentrotus purpuratus, puurchin, sperm genomic BAC library Strongylocentrotus genomic clone Plate=34 Col=12 Row=N, DNA sequence. AZ152840
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A2988471 642 bp DNA linear GSS 27 2M0271021F Mouse 10kb plasmid UUGC2M library Mus musculus clone UUGC2M0271021 F, DNA sequence.
A2988471
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California Institute of Technology
Pasadena California 91125, USA
Tel: (626) 395-8421
Fax: (626) 793-3047
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Class: BAC ends
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/clone=ib="Strongylocentrotus purpuratus, purple
/clone_lib="Strongylocentrotus purpuratus, purple
urchin_ sperm genomic BAC library"
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REFERENCE
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                                                                                                                  US-09-807-459-2 (1-458) x AZ988471 (1-642)
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Best Local Similarity:
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Mammalia; Eutheria; Rodentia;
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                                                                                                                                                                                                                                                                                                                                            10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114)gblAF129072.1), a copy-number inducible derivative of plasmid RI. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli xL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /lab_host="E. coli strain XL10-Gold, T1-resistant, /note="Vector: PWD42nv; Purified genomic DNA from musculus C57BL/6J (female) was obtained from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 adaptored DNA was purified and size-selected for a 9.5 to
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/clone_lib="Mouse 10kb plasmid UUGC2M library"
/sex="Female"
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/strain="C57BL/6J"
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                                                   Arakawa, T., Carninci, P., Fukuda, S., Furuno, M., Hanagaki, T., Hara, A., Hiramoto, K., Hori, F., Ishi, Y., Ito, M., Kawai, J., Konno, H., Kouda, M., Koya, S., Matsuyama, T., Miyazaki, A., Nomura, K., Ohno, M., Okazaki, Y., Okido, T., Saito, R., Sakai, C., Sakai, K., Sano, H., Sasaki, D., Shibata, K., Shinagawa, A., Shiraki, T., Sogabe, Y., Suzuki, H., Tagami, M., Tagawa, A., Takahashi, F., Takeda, Y., Tanaka, T., Toya, T., Muramatsu, M. and Hayashizaki, Y., Muramatsu, M. and Hayashizaki, Y., Muramatsu, M. and Hayashizaki, Y., Tanaka, T., et al. 2001)
Unpublished (2001)
On Jul 3, 2000 this sequence version replaced gi:8900669.
Contact: Yoshihide Hayashizaki
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BB230024
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                                Exploration Research Group,
   Institute
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URL:http://genome.gsc.riken.go.jp,
Carninci,P., Shibata,Y., Hayatsu,N., Sugahara,Y., Shibata,K., Itoh
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wagi,K., Fujiwake,S., Inoue,K., Togawa,Y., Izawa,M., Ohara,E.,
%atahiki,M., Yoneda,Y., Ishikawa,T., Ozawa,K., Tanaka,T., Matsuura
,S., Kawai,J., Okazaki,Y., Muramatsu,M., Inoue,Y., Kira,A. and
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Division of Experimental Animal Research in Riken contributed t
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Konno,H., Fukunishi,Y., Shibata,K., Itoh,M., Carninci,P., Sug,Y. and Hayashizaki,Y.
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Tel: 81-45-503-9222
Fax: 81-45-503-9216
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note-"Site_1: Sall; Site_2: BamHI; cDNA library was prepared and sequenced in Mouse Genome Encyclopedia Project of Genome Exploration Research Group in Riken Genomic Sciences Center and Genome Science Laboratory in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RIKEN. Division of Experimental Animal Research in Riken contributed to prepare mouse tissues. 1st strand cDNA was
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GCAGAAGATTCCATAGAGAAC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATCAAAGAAATGCAAAATGC----AAAAAGATCCTAACTCAA
                                 Tissue Procurement: Lothar Hennighausen Ph.D., Robin Hump
cDNA Library Preparation: Life Technologies, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                       Submitted (19-DEC-2001) National Institutes of Health, Mammalian Gene Collection (MGC), Cancer Genomics Office, National Cancer Institute, 31 Center Drive, Room 11A03, Bethesda, MD 20892-2590,
DNA Sequencing by: Baylor College of Medicine Human Genome Sequencing Center
                                                                                                      NIH-MGC Project URL: http://mgc.nci.nih.gov
Contact: MGC help desk
                                                                                                                                                                                                                                   Strausberg, R
                                                                                                                                                                                                                                                                                                       Mus musculus
                                                                                     Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                   Direct Submission
                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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Alignment
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Best Local Similarity:
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ValAsnArgValPheIleProThrLysLysPhePheAsnLysGluIleArgGluProSer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GluLysAsnValThrArgAspAlaAspAlaThrAspIleGluAsnPheAlaSerArgTyr 163
                                                                                   AlaGlyTyrValAspThrProTrpTyrLysLysTrpTyrMetLysLeuLysAsnPheMet
                                                                                                                                        ----AAGTCTCATAAGAAGACTAAAGAAGAAGTCGCAGGGACTCTGGAAGCAGTCCAA 458
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                                             GCCATTCAGAATATAACTCAGGCACTCCAGAAG
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This clone was selected for full length sequencing because it passed the following selection criteria: Hexamer frequency ORF analysis, GenomeScan gene prediction
This clone has the following problem: frame shifted.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Web site: http://www.hgsc.bcm.tmc.eau/c
Contact: amg@bcm.tmc.edu
Gunaratne, P.H., Garcla, A.M., Lu, X.,
Yoon, V.S., Kowis, C.R., Lawrence, S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Richards, S., Gibbs, R.A.
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/clone_lib="NCI_CGAP_Mam5"
/lab_host="DH10B"
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157 c 227 g 203 t
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/db_xref="taxon:10090"
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Martin, R.G., Muzny,
                                             -TCGAAGGAAAATTACACT
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У, D.M.,
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2M0082P07R Mouse 10kb plasmid UUGC1M library Mus musculus genomic
clone UUGC2M0082P07 R, DNA sequence.
                                                                                                                                                High
                                                                                                                                                                          Email: ddunn@genetlcs.utah.edu
Insert Length: 10000 Std Error: 0
Plate: 0082 row: P column: 07
Seg primer: CACACAGGAAACAGCTATGACC
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1 (bases 1 to 721)
Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
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University of Utah
Rm. 308, Biomedical
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Unpublished (2000)
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Fax: 801 585 7177
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/sex="Male"
                                                                                                                             Location/Qualifiers
                                              /db_xref="taxon:10090"
/clone="UUGC2M0082P07"
                                                                               /organism="Mus musculus"
/strain="C57BL/6J"
/lab_host="E.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Submitted (07-SEP-2000) Genoscope - Centre National de Sequencage, 2 rue Gaston Cremieux, CP 5706, 91057 EVRY cedex, FRANCE. (E-mail: seqreffgenoscope.cns.fr - Web: www.genoscope.cns.fr) This GSS is part of a random genomic sequencing program of thirteen yeast species: Saccharomyces bayanus var uvarum, Saccharomyces
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Souciet J.L., Aigle, M., Artiguenave, F., Blandin, G., Bolotin-Fukuhara, M., Bon, E., Brottier, P., Casaregola, S., de-Montigny, J., Dujon, B., Durrens, P., Lepingle, A., Llorente, B., Malpertuy, A., Neuveglise, C., Ozier-Kalogeropoulos, O., Potier, S., Saurin, W., Tekaia, F., Toffano-Nioche, C., Wesolowski-Louvel, M., Wincker, P. and Weissenbach, J. Genomic exploration of the hemiascomycetous yeasts: 1. A set of yeast species for molecular evolution studies
FEBS Lett. 487 (1), 3-12 (2000)
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Eukaryota; Fung1; Ascomycota; Saccharomycotina; Saccharomycetales;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
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                                                                complement(<4. .>939)
/note="similar to Saccharomyces cerevisiae ORF YFL008w
SMC1; chromosome segregation protein ]"
               /evidence=not_experimental
212 c 155 g 395 :
                                                                                                                                            /note="end : T3"
                                                                                                                                                                                             /db_xref="taxon:4931"
/clone="ASOAA012F08"
                                                                                                                                                                                                                                              /variety="uvarum"
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                                                                                                                                         PheIleAsnAsnGluIleArgAspProSerLysAlaLeuIleArgLysValTyrThrGlu 416
                                                                                                                                                                                                                                                                                                                               LysValSerThrGlyAlaGluAspLeuPheGluAsnLysIleGlyGlnGlyThrValAsp 396
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CNSO7C18
T3 end of clone xBC0AA00
of Debaryomyces hansenii
AL438594
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108.00
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                             hansenii, genomic survey sequence
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2 rue Gaston Cremieux, CP 5706, 91057 EPRY cedex - FRANCE (E-mail:
seqref@genoscope.cns.fr - Web: www.genoscope.cns.fr)
This GSS is part of a random genomic sequencing program of thirteen
yeast species: Saccharomyces bayanus var. uvarum, Saccharomyces
exiguus, Saccharomyces servazzii, Zygosaccharomyces rouxii,
Saccharomyces kluyveri, Kluyveromyces thermotolerans, Kluyveromyces
lactis var. lactis, Kluyveromyces marxianus var. marxianus, Pichia
angusta, Debaryomyces hansenii var. hansenii, Pichia sorbitophila,
Candida tropicalis and Yarrowia lipolytica. Genomic inserts of 3 to
5 kb were prepared and both extremities were sequenced. See
keywords for description of this sequence and for the sequence of
the other extremity of this insert.
AACGGATCG----
                                      ThrGlyLeuPheGlyTrpGlyIleLysArgAlaLeuLysGlnIleIleArgSerAsnLeu 211
                                                                                  AAGACTCTTCAGAGAATTGACGAA
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Soucitet, J.L., Aigle, M., Artiguenave, F., Blandin, G.,
Bolotin-Fukuhara, M., Bon, E., Brottier, P., Casaregola, S.,
Bolotin-Fukuhara, M., Durrens, P., Lepingle, A., Llorente, B.,
de-Montigny, J., Dujon, B., Durrens, P., Lepingle, A., Llorente, B.,
Malpertuy, A., Neuveglise, C., Ozier-Kalogeropoulos, O., Potier, S.,
Saurin, W., Tekala, F., Toffano-Nioche, C., Wesolowski-Louvel, M.,
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Debaryomyces hansenii
Eukaryota; Fungl; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Debaryomyces.
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                                                                                                                                                                                                                                                                                                                                                                                                                        268
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/note-"similar to Saccharomyces
weak similarity to spindle pole
/evidence-not_experimental
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/clone="XBCOAA001D03"
/clone_lib="XBCOAA"
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/strain="CBS 767"
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Mus musculus

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleoston

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleoston

Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; N

1 (bases 1 to 642)

Dunn, D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,

Junn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C.,

Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reil

,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,

and Wright,D., Weiss,R.
                                                                                                                                                                                          AZ431579 642 bp DNA linear IM0216006R Mouse 10kb plasmid UGGCIM library Mus clone UUGCIM0216006 R, DNA sequence.
AZ431579 AZ431579.1 GI:10555592
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                                    GluAsnLysIleGlyGlnGlyThrValAspPhePheAsnLysGluIleArgAspProSer 336
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                                                                                                                                                                    ATGATGGTAGAGGACATCAAGAAGAACTTTAATAACTCACTTAAAGAAATACAGGAGGAC
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Insert Length: 10000 Std Error:
Plate: 0216 row: O column: 06
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Fax: 801 585 7177
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University of Utah
Rm. 308, Biomedical
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Unpublished (2000)
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gh quality sequence stop:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (g114732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     USA
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/clone="UUGC1M0216006"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
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/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Shaying Zhao
Department of Eukaryotic Genomics
The Institute for Genomic Research
9712 Medical Center Dr., Rockville,
Tel: 301 838 0200
Fax: 301 838 0208
Email: szhao@tigr.org
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                                                                                                                                                                                                                                                                                                                   Clones are derived from the mouse BAC library RPCI-24. For BAC library availability, please contact Pieter de Jong (pdejong@mail.cho.org). Clones may be purchased from BACPAC Resources (http://www.chori.org/bacpac/orderingframe.htm). BAC page: http://www.tigr.org/tdb/bac_ends/mouse/bac_end_intro.html Plate: 149 row: N column: 17
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Tsegaye,G., Geer,K., Krol,M., Shvartsbeyn,A.,
Russell,D., de Jong,P. and Fraser,C.M.
Mouse BAC End Sequences from Library RPCI-24
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                                                                                    TTTGGTGTTCCAAGAGTGTTGACACAGGTTTTCAGA
                                                                                                                         TrpGlyIleLysArgAlaLeuLysGlnIleIleArgSerAsnLeuProLeuAspIleGly 216
                                     ThrGluHisSerValSerArgLeuGlnHisIleThrSerSerTyrLysAspTyrMetAsp 236
ACAAGCTCCAGCCAGAGTCAGTTATATCATCTAACACCAGAGATGACCAAATGGTGAAAG
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Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0
Plate: 0271 row: A column: 02
Seq primer: CACACAGGAAACAGCTATGACC
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Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   plasmid inserts
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,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
and Wright,D.,Weiss,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mouse whole genome scaffolding with
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pubMad 2 (g1|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid RI. The vector was ligated with adaptors complementary to the insert adaptors and continuous complementary to the insert adaptors and
                                                                                                                                                                                                                                                                                                                                                                                                 purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."

95 c 92 g 191 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGCIM0271A02"
/clone_lib="Mouse 10kb plasmid UUGCIM library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /lab_host="E. Coli strain XL10-Gold, T1-resistant,
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                                                                                                                                                                                                                                                                                                    Email: est@watson.wustl.edu
Library was constructed by R. Haywood. DNA sequencing by:
Washington University Genome Sequencing Center For information
obtaining a clone please contact: L. David Sibley
(sibley@borcim.wustl.edu), Washington University
                                                                                                                                                                                                                                                                                                                                                                                                             4444 Forest Park Parkway,
Tel: 314 286 1800
Fax: 314 286 1810
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Plasmodium falciparum 3D7.

---ium falciparum 3D7
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Eukaryota; Alveolata; Apicomplexa; Haemosporida; Plasmodium
1 (bases 1 to 532)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: L. David Sibley
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BM273998.1 GI:17967300
                                                                                                                                                                                                                                                            primer: -40UP from Gibco
h quality sequence stop: 422
/dev_stage="gametocyte (stage III-V)"
/lab_host="PH10B (GeneHog, Invitrogen, Inc.)"
/note="Vector: pBluescript SK plus; Site_1: EcoRI; Site_2:
XhoI; The library was constructed by R Haywood. cDNAs were synthesized from ganetocyte poly(A) + RNA by oligo d(T) priming, size-selected and directionally cloned into the
                                                                                                                                 library"
                                                                                                                                                       /clone_lib="Plasmodium
                                                                                                                                                                         /organism="Plasmodium falciparum
/db_xref="taxon:36329"
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Best Local Similarity:
                                                                                                                               SOURCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LeuLeuGluLys 458
                                                                                                                                                                                                                                                                                                                                                                                       GTAGATGAAGAAGTAACTGAAGAATTAATTGAAAAAGTAGATGAAGAAGTTGCTGAAGAA
                                                                                                                                                                                                                                                                                                                                                                                                                         GluIleArgAspProSerLysAlaLeuIleArgLysValSerThrGlu---AlaAspAsn 454
                                                                                                                                                                                                                                                                                                                                                                                                                                                             AlaAspAspLeuPheGluAsnLysIleGlyGlnGly---ThrValAspPheIleAsnLys
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAAGTAGATGAAGAAGTA-----GCTGAAGAATTAGTTGAAAAAGTAGATGAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PheIleAsnAsnGluIleArgAspProSerLysAlaLeuIleArgLysValTyrThrGlu 416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAGGTA-----GCTGATGAATTAGTTGAA---AAAGTAGCTGAAGAATTAGTTGAA
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 Dunn,D., Aoyagi,A., Barber,M., Beacorn,T.,
Islam,H., Longacre,S., Mahmoud,M., Meenen,I
,M., Rose,M., Rose,R., Stokes,R., Tingey,A
                                                                                                            Mus musculus
                                                                                                                                                  GSS.
                                                                                                                                                                 AZ944493
AZ944493.1 GI:13808782
                                                                                                                                                                                                   2M0205C06R Mouse 10kb plasmid UUGC2M clone UUGC2M0205C06 R, DNA sequence.
                                                                        Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia;
                                                     Mammalia; Eutheria;
1 (bases 1 to 596)
                                                                                                                               house mouse.
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                                                                                                                                                                                                                                                                                                                   462
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Matches:
Conservative:
Mismatches:
Indels:
                                                                    Craniata; Vertebrata;
Sciurognathi; Muridae
                                                                                                                                                                                                                    UUGC2M library Mus musculus
Tingey, A.,
                 Meenen, E.,
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                  Pedersen, T.,
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Best Local Similarity:
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                                                                                                                                                                                   243 ProLysPheAlaLysArgPheSerLeuMetValValGlnArgLeuLeuAlaThrValAla
                                                                                                          GlyTyrValAspThrProTrpTyrLysLysTrpTyrMetLysLeuLysAsnPheMetVal
                                                                                                                                                                                                                             AAACTTAAGAATCTTACTAACAGAAACCAAGACCACTCACCATCATCAGAACCCAGCACT
 ATGATGGTA----GAGGACATCAAGAAGGACTTTAATAACTCACTTAAAGAAATACCGGAG
                                 AsnArgValPheIleProThrLysLysPhePheAsnLysGluIleArgGlu----ProSer
                                                                         GGGCACCTCAACACCCC-----AAAAAGCTAGACCCAGATTTAAAAGCATATCTCATG
                                                                                                                                                  CCC----
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           University of Utah
University of Utah
Rm. 308, Biomedical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    plasmid inserts
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       quality sequence stop: 596
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  92
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114|gb|AF129072.1), a copy number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XIIO-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /lab_host="E. coli strain XL10-Gold, T1-resistant, /note="Vector: PWD42nv; Purified genomic DNA from I musculus C57BL/6J (female) was obtained from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Biomedical Polymers Research
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /db_xref="taxon:10090"
/clone="UUGC2M0205C06"
/clone_lib="Mouse 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Mus musculus'
/strain="C57BL/6J"
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Plasmodium yoelii yoelii.

Plasmodium yoelii yoelii

Eukaryota; Alveolata; Apicomplexa; Haemosporida;
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                                                                                                                                                                                                                                                                                                                                                                                                                           Parasite Genomics Group
The Institute for Genomic Research
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                                                                                                                                                                                                                                                                                                      http://www.malaria.mr4.org/mr4pages/index.html
                                                                                                                                                                                                                                                                                                                        For clone info, please contact the Malaria Reagent Resource Center, ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Jane Carlton
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          isolated using the guanidinium isothiocyanate method, a mRNA isolated using oligo(dT)-cellulose chromatography. First strand cDNA synthesis was completed using a 50-ba
                                                               /note="Vector: pAD-GAL4; At 20-25% parasitemia, blocollected from BALB/CByJ mice infected with Py17XL parasites, and leukocytes removed by passage over microcrystalline cellulose collumns. Total RNA was
primer and reverse transcriptase in the presence
                                                                                                                                /dev_stage="Asexual blood stages"
/lab_host="E. coli XL-1 Blue"
                                                                                                                                                                                    /clone="PYCPA68"
                                                                                                                                                                    /clone_lib="PyBS"
                                                                                                                                                                                                       /db_xref="taxon:73239"
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RESULT 28
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                                                                                          DNA sequence.
BH126086
                                                                                                                                  RPCI-24-289K13.TJ RPCI-24 Mus
                                                                                                                                                      вн126086
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                 Mus musculus
                                                                           BH126086.1 GI:14969598
                                     house mouse.
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                              ThraspThrLysAspLeuPheGlu---AsnLysIleGlyGlnGlyThrValAspPhePhe 328
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                                                                                                                                                                                                                                                                                         AspThrGlnIleProAlaLeuProLysPheAlaLysArgPheSerLeuMetValValGln
                                                                                                                                                                                                                                                            library availability, please contact Pieter de Jong (pdejong@mail.cho.org). Clones may be purchased from BACPAC Resources (http://www.chori.org/bacpac/orderingframe.htm). BAC page: http://www.tigr.org/tdb/bac_ends/mouse/bac_end_intro.html plate: 289 row: K column: 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Department of Eukaryotic Genomics
The Institute for Genomic Research
9712 Medical Center Dr., Rockville,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Zhao, S., Nierman, W., Malek, J., Shatsman, S., Ak
Tsegaye, G., Geer, K., Krol, M., Shvartsbeyn, A.,
Russell, D., de Jong, P. and Fraser, C.M.
Mouse BAC End Sequences from Library RPCI-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unpublished (1999)
Other_GSSs: RPCI-24-289K13.TV
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/cell_type="Spleen/Brain"
/note="Vector: pTARBAC1; Site_1: BamH1; Site_2: BamH1;
/note="Vector: pTARBAC1; Site_1: BamH1; Site_2: BamH1;
/note="Vector pTARBAC1; Site_1: BamH1; Site_2: BamH1;
/note="Vector pTARBAC1; Site_2: BamH1; Site_2: BamH1; Sites using MboI partially digested male C57BL/6J
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/clone="RPCI-24-289K13"
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Contact: Jim Brandle
Genomics and Biotechnology
Agriculture and Agri-Food Canada -
1391 Sandford St., London, Ontario,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Leaf ESTs from Stevia rebaudiana: diterpene synthesis
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Brandle, J.E., Richman, A.,
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519 457 1470
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/tissue_type="leaf"
/tissue_type="leaf"
/dev_stage="field grown, mid-size"
/dev_stage="field grown, mid-size"
/lab_host="E. coli strain XLOLE"
/lab_host="E.
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                                                                                                                                                                                                                     products were digested with ECOKI and AIDLE ELEVATORY directional cloning into the lambda ZAP Express vector. The library was amplified using the host strain XLI-Blue MRF'. Mass excision of the library was performed to
                                                                                             obtain pBK-CMV phagemid clones in the host strain XLOLR. Single pass DNA sequencing was performed using the T3 promoter primer: 5' ATTAACCCTCACTAAAGGGA 3'. This library
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/strain="751/1501"
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                                                         Alex Richman."
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Zhao,S., Nierman,W., Malek,J., Shatsman,S., A) Tsegaye,G., Geer,K., Krol,M., Shvartsbeyn,A., Russell,D., de Jong,P. and Fraser,C.M.
Mouse BAC End Sequences from Library RPCI-24 Unpublished (1999)
Other,GSSs: RPCI-24-242M13.TV
Contact: Shaying Zhao
                                                                                     Mus musculus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;

1 (bases 1 to 791)
                                                                                                                                                                                                                 RPCI-24-242M13.TJ RPCI-24 Mus musculus genomic clone RPCI-24-242M13
                                                                                                                                                                             BH047254.1
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Percent Similarity:
Best Local Similarity:
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                                           ArgAspProSerLysAlaLeuLys···--GluLysValSerAsnAspAlaLysAspLeu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: szhao@tigr.org
Clones are derived from the mouse BAC library RPCI-24. For BAC library availability, please contact Pieter de Jong (pdejong@mail.cho.org). Clones may be purchased from BACPAC Resources (http://www.chori.org/bacpac/orderingframe.htm). BAC page: http://www.tigr.org/tdb/bac_ends/mouse/bac_end_intro.html Plate: 242 row: M column: 13
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The Institute for Genomic Research
9712 Medical Center Dr., Rockville.
Tel: 301 838 0200
Fax: 301 838 0208
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/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="RPCI-24-242M13"
/clone_lib="RPCI-24"
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/cell_type="Spleen/Brain"
/cell_type="Spleen/Brain"
/note="Vector: pTARBAC1; Site_1: BamH1; Site_2: BamH1;
/note="Vector: pTARBAC1; Site_1: BamH1; Site_2: BamH1;
RPCI-24 Mouse BAC Library produced by Pieter de Jong. The
RPCI-24 Mouse BAC Library produced by Pieter de Jong. The
library was cloned in the pTARBAC1 cloning vector at the
BamH1 sites using MboI partially digested male C57BL/6J
DNA."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      144 AAACCTATGGATAATAGGAATTGATGACACTGAAGATTTT------CAACTTAAAGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             aLeuIleArgLysValTyrThr------GluAlaAspAspLeuPheGluAsnLy 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nLysIleGlyGlnGlyThrValAspPheIleAsnAsnGluIleArgAspProSerLysAl 408
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ACCAGCAAATATCTTCAACAGAATTATAGAAGAAAACTTCCCAAACCTA 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       pProSerLysAlaLeuIleArgLysValSerThrGluAlaAspAsnLeu 455
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AGAGATGGAAGAGAATC----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gAspProSerLysAlaLeuIleArgLysValSerThrGlyAlaGluAspLeuPheGluAs 388
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Rei,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern
                                                                                                                                                                                                                                                                                                                                                                 Seq primer: CGTTGTAAAACGACGGCCAGTClass: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                        Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0208 row: J column: 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (2000)
Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                plasmid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mouse whole genome scaffolding with paired
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and Wright, D., Weiss, R.
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Mammalia; Eutheria; Rodentia;
1 (bases 1 to 439)
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2M0208J13F Mouse 10kb plasmid UUGC2M
clone UUGC2M0208J13 F, DNA sequence.
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801 585 7177
                                                                                                                                                                                                                                                                                                                    quality sequence stop: 439
Location/Qualifiers
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                 (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA
                                                                                                   /lab_host="E. coli strain XL10-Gold, T1-resistant, /note="Vector: PWD42nv; Purified genomic DNA from I musculus C57BL/6J (female) was obtained from the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Biomedical
                                                                                      Laboratory Mouse DNA Resource
                                                                                                                                                                                                               /db_xref-"taxon:10090"
/clone-"UUGC2M0208J13"
                                                                                                                                                                                                                                                         /organism="Mus musculus"
/strain="C57BL/6J"
                                                                                                                                                                                         /clone_lib-"Mouse 10kb plasmid UUGC2M library"
                                                                                                                                                                         /sex="Female"
blunt end-repaired
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VERSION
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Best Local Similarity:
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                                                                  REFERENCE
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                                                                                                                                                                                                                                                                                                                                                      98
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               LysValSerThrGlyAlaGluAspLeuPheGluAsnLysIleGlyGlnGlyThrValAsp
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CAAGTCTTTAAAGAAAAACAGGAAAACACATTCAAACAGGTGATGGAA-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ---- AAGAAGGACTTTAATAACTTACTTAAAGAAATACAGAACACTGGTAAAGAGTTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ProThrLysLysPhePheAsn------LysGluIleArgGluProSerLysAlaLeu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GAAAAGTTAGAACTGGAGTTAAAAACATATCACATGATGATGGTAGAGGACATC----
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                                                                                                                                                                                                                                                                                                                                                                                PheIleAsnAsnGluIleArgAspProSerLysAlaLeuIleArgLysValTyrThrGlu 416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGAACCATAGATGCGAGCATCAGCAACAGAATACAAGAGATGGAAGAGAGAATC -----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GlyThrValAsp---PheIleAsnAsnGluIleArgAspProSerLysAlaLeuIleArg 376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -----LysGluLysValSerThrAspThrLysAspLeuPheGluAsnLysIleGly
Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., D
Islam,H., Longacre,S., Mahmoud,M., Meenen,E.,
,M., Rose,M., Rose,R., Stokes,R., Tingey,A.,
                                                                                                                                                                                                                             AZ344512 503 bp DNA li
1M0078M05R Mouse 10kb plasmid UUGC1M library
clone UUGC1M0078M05 R, DNA sequence.
                                                                                                                           Mus musculus
                                                                                                                                                                      GSS
                                                                                   Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                                                                                                                nouse mouse
                                                                                                                                                                                                                                                                                                                                                        ATCGGCAAAACAATCAAAGAA----AAGGCAAAATGCAAAAAGATCTTAACTCAG
                                                         ummalia; Eutheria; (bases 1 to 503)
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                                                                                                                                                                                                                                                                                                                                                                                                                                       TCAGGTGCAGAAGATTCCATAGAGAAC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Strand and selected for amplicillin resistance."

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ligated to the blunt ends in high molar excess. The
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30.00%
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                                                                                   Chordata;
Rodentia;
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Matches:
Conservative:
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                                                                                Craniata; Veri
Sciurognathi;
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Meenen, E., rc.
                                                                                                        Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     439
48
27
40
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                       Duval,B., Hamil,C
E., Pedersen,T., Re
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                                                                                     Muridae;
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                                                                                     Murinae;
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                                                                                                                                                                412
340
                                         283
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                                                                                                                                                                                                        243
                                                                                                                                                                                                                                               472 AAATGTGAGAATCTTACTAACAGAAAACAGGACCACTCACCATCATCAGAACCCAGCATT
ATGATGGTA---GAGGACCAAGAAGGACTTAAATAACTCACTTAAAGAAATACAGGAG
                 AsnArgValPheIleProThrLysLysPhePheAsn------LysGluIleArgGlu
                                                                              GGGCACCCCAACACACCCC----AAAAAGCTAGACCTGGATTTAAAAAGCATATCTCATG
                                                                                                        GlyTyrValAspThrProTrpTyrLysLysTrpTyrMetLysLeuLysAsnPheMetVal
                                                                                                                                                                                                     ProLysPheAlaLysArgPheSerLeuMetValValGlnArgLeuLeuAlaThrValAla
                                                                                                                                                                                                                                                                                      ArgLeuGlnHisIleThrSerSerTyrLysAspTyrMetAspThrGlnIleProAlaLeu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 University of Utah Genome Center University of Utah Rm. 308, Biomedical Polymers Res
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Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0078 row: M column: 05
Seq primer: CACACAGGAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and Wright,D.,Weiss,R. Mouse whole genome scaffolding with paired end reads from 10 \, \mathrm{kb}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with 74 DNA polymerase and 74 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pMTA2 (gil4732114|gblAF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note-"Vector: PWD42nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0078M05"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Email: szhao@tigr.org
Clones are derived from the mouse BAC library RPCI-23. For BAC
Clones availability, please contact Pieter de Jong
(pdejong@mail.cho.org). Clones may be purchased from BACPAC
Resources (http://www.chori.org/bacpac/orderingframe.htm). BAC
page: http://www.tigr.org/tdb/bac_ends/mouse/bac_end_intro.htm
Plate: 214 row: L column: 9
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Fax: 301 838 0208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Department of Eukaryotic Genomics
The Institute for Genomic Researc
9712 Medical Center Dr., Rockvill
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GSS.
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,B., Levins,M., Mcgann,S., Tsegaye,G., Geer,K., Krol,M., de Jong,P
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
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                                                                                                  Contact: Brewus.
Department of Eukaryotic Genomics
The Institute for Genomic Research
The Institute for Genomic Rockville,
 DNA
                                                                        9712 Medical Center Dr., Tel: 301 838 0208 Fax: 301 838 3543
                                    Email: bjloftus@tigr.org Clones are derived from
                                                                                                                                                                                  Unpublished (2000)
                                                                                                                                                                                                   Loftus, B., Van Aken, S. and Fraser, C. Determination of clone end sequences HM1: IMSS sheared DNA library
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Eukaryota; Entamoebidae;
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GSS.
Entamoeba histolytica.
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                                                                                       IleGlyGlnGlyThrValAspPhePheAsnLysGluIleArgAspProSerLysAlaLeu
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quality sequence stop: {
Location/Qualifiers
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/Clone_Tyector: pHOS1; Site_1: Bst I; Constructed at The
/note="Yector: pHOS1; Site_1: Bst I; Constructed at The
/note="Yector: pHOS1; Site_1: Constructed at The
/note="Yector: pHOS1; Site_1: Associated from broth cultures of E. histolytica
Genomic DNA isolated from broth cultures of E. histolytica
using a method described by Clark and Diamond (Clark,
C.G., and Diamond, L.S. (1993) Entamoeba histolytica: a
method for isolate identification. Exp. Parasitol.
77:450.) The DNA was mechanically sheared to give a
tight size distribution (-2 kb). The v + i method used for
the library construction is described in detail in Smith,
H.O. and Venter, J.C. (Making small insert libraries for
whole genome shotgun sequencing projects. In Genome
Sequencing: A Practical Approach, eds. M. Vaudin and B.
Barell, Oxford University Press, 1999)."
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/strain-"HM1:IMSS"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: Szhao@tigr.org
Clones are derived from the mouse BAC library RPCI-24. For BAC
Clones availability, please contact Pieter de Jong
(pdejong@mail.cho.org). Clones may be purchased from BACPAC
Resources (http://www.chori.org/bacpac/orderingframe.htm). BAC page: http://www.tigr.org/tdb/bac_ends/mouse/bac_end_intro.html
Plate: 275 row: B column: 18
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Department of Eukaryotic Genomics
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Other_GSSs: RPCI-24-275B18.TJ
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Tsegaye,G., Geer,K., KrOl,M., Shvartsbeyn,A.,
Russell,D., de Jong,P. and Fraser,C.M.
Mouse BAC End Sequences from Library RPCI-24
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Mammalia; Eutheria; Rodentia;
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                                                                                                                                                                                                                                           /cell_type="Spleen/Brain" /cell_type="Spleen/Brain" /note="Vector: pTARBAC1; Site_1: BamH1; Site_2: BamH1; /note="Vector: pTARBAC1; Site_1: BamH1; Site_2: BamH1; /note="Vector by Pleter de Jong. The library was cloned in the pTARBAC1 cloning vector at the BamH1 sites using MboI partially digested male C57BL/6J DNA."
                                                                                                                                                                                                                                                                                                                                                    /db_xref="taxon:10090"
/clone="RPCI-24-275B18"
/clone_lib="RPCI-24"
                                                                                                                                                                                                                                                                                                                                     /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                   /organism="Mus musculus"
/strain="C57BL/6J"
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             9712 Medical Center Dr., Tel: 301 838 0208 Fax: 301 838 3543
                                                                                                                                                                                                                                                     genomic, DNA sequence.
AZ667704
AZ667704.1 GI:11804970
                                                            Department of Eukaryotic Genomics The Institute for Genomic Research
                                                                                                                        Eukaryota; Entamoebidae; Entamoeba.

1 (bases 1 to 830)

Loftus, B., Van Aken, S. and Fraser, C. Determination of clone end sequences

HM1: IMSS sheared DNA library
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                                                                                          Contact: Brendan J Loftus
                                                                                                         Unpublished (2000)
Email: bjloftus@tigr.org
                                                                                                                                                                                                         Entamoeba histolytica
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Entamoeba histolytica
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DB:
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Best Local Similarity:
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Seq primer: M13-Forward
Class: shotgun
High quality sequence start: 33
High quality sequence stop: 815.
Location/Qualifiers
830
      ATAÇATGTGATAGAAGGATACTTAGAT-----
                                                   LeuAlaThrValAlaGlyTyrValAspThrProTrpTyrLysLysTrpTyrMetLysLeu 277
                                                                                                                                                                                                                                                                                                                                                                                                         GlyIleLysArgAlaLeuLysGlnIleIleArgSerAsnLeuProLeuAspIleGlyThr 217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LeuTyrMetAlaThrLeu-----TyrTyrLysThrTyrThrAsnValAspGluPheGly 181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GluLysAsnValThrArgAspAlaAspAlaThrAspIleGluAsnPheAlaSerArgTyr 163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MetArgPheArgArgGlyLysAsnHisSerTyrPheHisAspLeuValPheAsnLeuLeu 143
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                                                                                                                     TCACAAGTAATTTTATTTACATTCCAAAAGAAATAT-----
                                                                                                                                                                        GlnIleProAlaLeuProLysPheAlaLysArgPheSerLeuMetValValGlnArgLeu 257
                                                                                                                                                                                                                                       CGATCAGAAATTGAAAGTTTAGTCCAT-----
                                                                                                                                                                                                                                                                                       GluHisSerValSerArgLeuGlnHisIleThrSerSerTyrLysAspTyrMetAspThr 237
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /strain="HMI:IMSS"
/db_xref="taxon:9759"
/clone_lib="Entamoeba histolytica Sheared DNA"
/clone_lib="Entamoeba histolytica Sheared DNA"
/clone_lib="Entamoeba histolytica Structed at The
/note="yector: pHOS1; Site_1: Bst I; Constructed at The
/note="yector: pHOS1; Site_1: 
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Matches:
Conservative:
Mismatches:
Indels:
Gaps:
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FEATURES Sou	TIT JOU COMME	REFER AUT	ACCESSION VERSION KEYWORDS SOURCE ORGANIS	RESUL AZ891 LOCUS DEFIN	β 6γ	Db .	g b	γΩ	Db Qy	Ωу	ру	ОУ	Qy Db
	TITLE JOURNAL COMMENT	EFERENCE AUTHORS	CESSION RSION YWORDS URCE ORGANISM	RESULT 37 AZ891746/c LOCUS DEFINITION	400		159	366	358 219	338 261		298 339	
Department of Eukaryotic Genomics The Institute for Genomic Research 9712 Medical Center Dr., Rockville, MD 20850, USA Tel: 301 838 0200 Fax: 301 838 0208 Email: szhao@tigr.org Clones are derived from the mouse BAC library RPCI-24. For BAC library availability, please contact Pieter de Jong (pdejong@mail.cho.org). Clones may be purchased from BACPAC Resources (http://www.chori.org/Dacpac/orderingframe.htm). BAC end page: http://www.tigr.org/tdb/bac_ends/mouse/bac_end_intro.html plate: 209 row: J column: 3 Seq primer: T7 Class: BAC ends. Location/Qualifiers 1605 /organism="Mus musculus" /strain="C57BL/6J" /db_xref="taxon:10090" /clone="RPCI-24-209J3"	er,C.M. brary RPCI-24	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 605) Zhao, S., Nierman, W., Malek, J., Shatsman, S., Akinret, B., Levins, M., Tsedave, G., Geer, K., Krol, M., Shvartsbevn, A., Gebredeorgis, E.,	A2891746 Az891746.1 GI:13210691 GSS. house mouse. Mus musculus	AZ891746 605 bp DNA linear GSS 05-MAR-2001 RPCI-24-209J3.TV RPCI-24 Mus musculus genomic clone RPCI-24-209J3, DNA sequence.	AsnGluIleArgAspProSerLysAlaLeuIleArgLysVal 413 :::::	TGTTATCAAACATTTATTACAAAATCATCAATACAATTTGAATTTATAACA	CAAGAAGATCAAGAAATAACAGATATTATTTCTATTATTTCA 118		GlyThrValAspPheIleAsnAsn	spAlaLysAspLeuPheGluAsnLysILeGlyGln ::: 		ArgGluProSerLysAlaLeuLysGluLysValSerThrAspThrLysAspLeuPheGlu 317 ::: :::	le AT

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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus

1 (bases 1 to 614)
                                                                                                                                                                                                                                                                                                             AZ415896 614 bp DNA linear GSS UJ-UCT-ZUU 1M0190A17R Mouse 10kb plasmid UUGC1M library Mus musculus genomic clore.
Unpublished (2000)
                    plasmid inserts
                                    Mouse whole genome scaffolding with paired end reads from 10kb
                                                          Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A., and Wright, D., Weiss, R.
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/Cell_Type="Spleen/Brain"
/Cell_Type="Spleen/Brain"
/Cell_Type="Spleen/Brain"
/note="Vector: pTARBAC1; Site_1: BamH1; Site_2: BamH1;
/note="Vector: pTARBAC1; Site_1: BamH1; Site_2: BamH1;
/note="Vector: pTARBAC1; Site_2: BamH1;
/note="Vector: pTARBAC1; Site_2: BamH1; Sites using MboI partially digested male C57BL/6JDNA."
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                                                                                   GTAGAGGACATCAAGAAGGACTTAAATAACTCACTTAAAGAAATACAGGAGAACACTGCT
                                                                                                                                                                                                              ValAspThrProTrpTyrLysLysTrpTyrMetLysLeuLysAsnPheMetValAsnArg
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                                   -----LysAlaLeuLysGluLysValSerThrAspThrLysAspLeuPheGluAsn
                                                                                                                           ValPheIleProThrLysLysPhePheAsn-----LysGluIleArgGluProSer---
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University of Utah G
University of Utah
Rm. 308, Biomedical
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Insert Length: 10000 Std Err
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Fax: 801 585 7177
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/clone="UUGC1M0190A17"
/clone_lib="Mouse 10kb plasmid UUGC1M library"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Mus musculus"
/strain="C57BL/6J"
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/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jack
Laboratory Mouse DNA Resource
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(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA
                                                                                                                                                                                                                          /organism="Mus musculus"
/strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="UUGC1M0465N20"
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BH313240 700 bp DNA
CH230-101L12.TV CHORI-230 Segment 1 Rattus
CH230-101L12, DNA sequence.
BH313240
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ArgAspProSerLysAlaLeuIleArgLysValSerThrGlyAlaGluAspLeuPheGlu 387 :::::: caaGaGaTGGAAGAGAGATCTCAGGAGCAGAAGAT 243	ASPLeuPheGluAsnLySIleGlyGlnGlyThrValAspPheIleAsnAsnGluIle 367 :::	CAAAAATCCCTGAAAGAATTCCAGGAAAACACAATCAAACAG	459-2 (1-458) x BH313240 (1-700) ThrLysLysPhePheAsnLysGluIleArgGluProSerLysAlaLeuLysGluLysVal 308 ::::: ::::: ACACAGGAAAACATAAATAAACAAGTA	res arí	Bukaryota: Metaz Mammalia; Euther Rattus. 1 (bases 1 to 7 Zhao, S., Shetty, , A., Gebregeorgi Jong, P. and Fras Rat BAC End Sequ Unpublished (199 Other_GSSs: CH23 Contact: Shaying Department of Eu The Institute fo 9712 Medical Cen Tel: 301 838 020 Email: szhaoetig Clones are deriv (http://www.chor page: http://www.chor page: http://www.page: http://www Plate: 101 row: Seq primer: F7 Class: BAC ends. Locatio 1. 7000 /organi /strain /db_xre /clone= /clone= /sex-"F /cell_t /note="" CHORI-2 Ppieter: 133 113 a 133	

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